

# LISTE DE SEQUENCES

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<120> NUCLEIC ACIDS OF THE HUMAN ABCA5, ABCA6, ABCA9, ABCA10 GENES,  
VECTORS CONTAINING SUCH NUCLEIC ACIDS AND USES THEREOF

<130> ABCA5, 6, 9, 10

<140> 03806.0529-00000

<141> 2001-12-07

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<170> PatentIn Ver. 2.1

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1630

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Tyr Ala Ala Ser Leu Val Phe Phe Ile Tyr Met Ile Ser Phe Ile Phe		
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1185	1190	1195
Phe Ile Pro Tyr Phe Gln Thr Leu Leu Phe Val Phe Val Leu Arg Cys		
1205	1210	1215
Met Glu Leu Lys Cys Gly Lys Lys Arg Met Arg Lys Asp Pro Val Phe		
1220	1225	1230
Arg Ile Ser Pro Gln Ser Arg Asp Ala Lys Pro Asn Pro Glu Glu Pro		
1235	1240	1245
Ile Asp Glu Asp Glu Asp Ile Gln Thr Glu Arg Ile Arg Thr Ala Thr		
1250	1255	1260
Ala Leu Thr Thr Ser Ile Leu Asp Glu Lys Pro Val Ile Ile Ala Ser		
1265	1270	1275
Cys Leu His Lys Glu Tyr Ala Gly Gln Lys Lys Ser Cys Phe Ser Lys		
1285	1290	1295
Arg Lys Lys Lys Ile Ala Ala Arg Asn Ile Ser Phe Cys Val Gln Glu		
1300	1305	1310
Gly Glu Ile Leu Gly Leu Leu Gly Pro Ser Gly Ala Gly Lys Ser Ser		
1315	1320	1325
Ser Ile Arg Met Ile Ser Gly Ile Thr Lys Pro Thr Ala Gly Glu Val		
1330	1335	1340

Glu Leu Lys Gly Cys Ser Ser Val Leu Gly His Leu Gly Tyr Cys Pro  
1345 1350 1355 1360

Gln Glu Asn Val Leu Trp Pro Met Leu Thr Leu Arg Glu His Leu Glu  
1365 1370 1375

Val Tyr Ala Ala Val Lys Gly Leu Arg Lys Ala Asp Ala Arg Leu Ala  
1380 1385 1390

Ile Ala Arg Leu Val Ser Ala Phe Lys Leu His Glu Gln Leu Asn Val  
1395 1400 1405

Pro Val Gln Lys Leu Thr Ala Gly Ile Thr Arg Lys Leu Cys Phe Val  
1410 1415 1420

Leu Ser Leu Leu Gly Asn Ser Pro Val Leu Leu Leu Asp Glu Pro Ser  
1425 1430 1435 1440

Thr Gly Ile Asp Pro Thr Gly Gln Gln Gln Met Trp Gln Ala Ile Gln  
1445 1450 1455

Ala Val Val Lys Asn Thr Glu Arg Gly Val Leu Leu Thr Thr His Asn  
1460 1465 1470

Leu Ala Glu Ala Glu Ala Leu Cys Asp Arg Val Ala Ile Met Val Ser  
1475 1480 1485

Gly Arg Leu Arg Cys Ile Gly Ser Ile Gln His Leu Lys Asn Lys Leu  
1490 1495 1500

Gly Lys Asp Tyr Ile Leu Glu Leu Lys Val Lys Glu Thr Ser Gln Val  
1505 1510 1515 1520

Thr Leu Val His Thr Glu Ile Leu Lys Leu Phe Pro Gln Ala Ala Gly  
1525 1530 1535

Gln Glu Arg Tyr Ser Ser Leu Leu Thr Tyr Lys Leu Pro Val Ala Asp  
1540 1545 1550

Val Tyr Pro Leu Ser Gln Thr Phe His Lys Leu Glu Ala Val Lys His  
1555 1560 1565

Asn Phe Asn Leu Glu Glu Tyr Ser Leu Ser Gln Cys Thr Leu Glu Lys  
1570 1575 1580

Val Phe Leu Glu Leu Ser Lys Glu Gln Glu Val Gly Asn Phe Asp Glu  
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Glu Ile Asp Thr Thr Met Arg Trp Lys Leu Leu Pro His Ser Asp Glu  
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Pro

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 <212> PRT  
 <213> Homo sapiens

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 Glu Trp Leu Phe Ser Phe Leu Leu Val Leu Phe Leu Tyr Leu Phe Phe  
 35 40 45  
 Ser Asn Leu His Gln Val His Asp Thr Pro Gln Met Ser Ser Met Asp  
 50 55 60  
 Leu Gly Arg Val Asp Ser Phe Asn Asp Thr Asn Tyr Val Ile Ala Phe  
 65 70 75 80  
 Ala Pro Glu Ser Lys Thr Thr Gln Glu Ile Met Asn Lys Val Ala Ser  
 85 90 95  
 Ala Pro Phe Leu Lys Gly Arg Thr Ile Met Gly Trp Pro Asp Glu Lys  
 100 105 110  
 Ser Met Asp Glu Leu Asp Leu Asn Tyr Ser Ile Asp Ala Val Arg Val  
 115 120 125  
 Ile Phe Thr Asp Thr Phe Ser Tyr His Leu Lys Phe Ser Trp Gly His  
 130 135 140  
 Arg Ile Pro Met Met Lys Glu His Arg Asp His Ser Ala His Cys Gln  
 145 150 155 160  
 Ala Val Asn Glu Lys Met Lys Cys Glu Gly Ser Glu Phe Trp Glu Lys  
 165 170 175  
 Gly Phe Val Ala Phe Gln Ala Ala Ile Asn Ala Ala Ile Ile Glu Ile  
 180 185 190  
 Ala Thr Asn His Ser Val Met Glu Gln Leu Met Ser Val Thr Gly Val  
 195 200 205  
 His Met Lys Ile Leu Pro Phe Val Ala Gln Gly Gly Val Ala Thr Asp  
 210 215 220  
 Phe Phe Ile Phe Phe Cys Ile Ile Ser Phe Ser Thr Phe Ile Tyr Tyr  
 225 230 235 240

Val	Ser	Val	Asn	Val	Thr	Gln	Glu	Arg	Gln	Tyr	Ile	Thr	Ser	Leu	Met
				245					250					255	
Thr	Met	Met	Gly	Leu	Arg	Glu	Ser	Ala	Phe	Trp	Leu	Ser	Trp	Gly	Leu
			260					265					270		
Met	Tyr	Ala	Gly	Phe	Ile	Leu	Ile	Met	Ala	Thr	Leu	Met	Ala	Leu	Ile
		275					280					285			
Val	Lys	Ser	Ala	Gln	Ile	Val	Val	Leu	Thr	Gly	Phe	Val	Met	Val	Phe
	290					295					300				
Thr	Leu	Phe	Leu	Leu	Tyr	Gly	Leu	Ser	Leu	Ile	Thr	Leu	Ala	Phe	Leu
305					310					315					320
Met	Ser	Val	Leu	Ile	Lys	Lys	Pro	Phe	Leu	Thr	Gly	Leu	Val	Val	Phe
				325					330					335	
Leu	Leu	Ile	Val	Phe	Trp	Gly	Ile	Leu	Gly	Phe	Pro	Ala	Leu	Tyr	Thr
			340					345					350		
His	Leu	Pro	Ala	Phe	Leu	Glu	Trp	Thr	Leu	Cys	Leu	Leu	Ser	Pro	Phe
		355					360					365			
Ala	Phe	Thr	Val	Gly	Met	Ala	Gln	Leu	Ile	His	Leu	Asp	Tyr	Asp	Val
	370					375					380				
Asn	Ser	Asn	Ala	His	Leu	Asp	Ser	Ser	Gln	Asn	Pro	Tyr	Leu	Ile	Ile
385					390					395					400
Ala	Thr	Leu	Phe	Met	Leu	Val	Phe	Asp	Thr	Leu	Leu	Tyr	Leu	Val	Leu
				405					410					415	
Thr	Leu	Tyr	Phe	Asp	Lys	Ile	Leu	Pro	Ala	Glu	Tyr	Gly	His	Arg	Cys
			420					425					430		
Ser	Pro	Leu	Phe	Phe	Leu	Lys	Ser	Cys	Phe	Trp	Phe	Gln	His	Gly	Arg
		435					440					445			
Ala	Asn	His	Val	Val	Leu	Glu	Asn	Glu	Thr	Asp	Ser	Asp	Pro	Thr	Pro
	450					455					460				
Asn	Asp	Cys	Phe	Glu	Pro	Val	Ser	Pro	Glu	Phe	Cys	Gly	Lys	Glu	Ala
465					470					475					480
Ile	Arg	Ile	Lys	Asn	Leu	Lys	Lys	Glu	Tyr	Ala	Gly	Lys	Cys	Glu	Arg
				485					490					495	
Val	Glu	Ala	Leu	Lys	Gly	Val	Val	Phe	Asp	Ile	Tyr	Glu	Gly	Gln	Ile
			500					505					510		
Thr	Ala	Leu	Leu	Gly	His	Ser	Gly	Ala	Gly	Lys	Thr	Thr	Leu	Leu	Asn
								30							

	515					520					525				
Ile	Leu 530	Ser	Gly	Leu	Ser	Val 535	Pro	Thr	Ser	Gly	Ser 540	Val	Thr	Val	Tyr
Asn 545	His	Thr	Leu	Ser	Arg 550	Met	Ala	Asp	Ile	Glu 555	Asn	Ile	Ser	Lys	Phe 560
Thr	Gly	Phe	Cys	Pro 565	Gln	Ser	Asn	Val	Gln 570	Phe	Gly	Phe	Leu	Thr 575	Val
Lys	Glu	Asn	Leu 580	Arg	Leu	Phe	Ala	Lys 585	Ile	Lys	Gly	Ile	Leu 590	Pro	His
Glu	Val	Glu 595	Lys	Glu	Val	Gln	Arg 600	Val	Val	Gln	Glu	Leu 605	Glu	Met	Glu
Asn	Ile 610	Gln	Asp	Ile	Leu	Ala 615	Gln	Asn	Leu	Ser	Gly 620	Gly	Gln	Asn	Arg
Lys 625	Leu	Thr	Phe	Gly	Ile 630	Ala	Ile	Leu	Gly	Asp 635	Pro	Gln	Val	Leu	Leu 640
Leu	Asp	Glu	Pro	Thr 645	Ala	Gly	Leu	Asp	Pro 650	Leu	Ser	Arg	His	Arg 655	Ile
Trp	Asn	Leu	Leu 660	Lys	Glu	Gly	Lys	Ser 665	Asp	Arg	Val	Ile	Leu 670	Phe	Ser
Thr	Gln	Phe 675	Ile	Asp	Glu	Ala	Asp 680	Ile	Leu	Ala	Asp	Arg 685	Lys	Val	Phe
Ile	Ser 690	Asn	Gly	Lys	Leu	Lys 695	Cys	Ala	Gly	Ser	Ser 700	Leu	Phe	Leu	Lys
Lys 705	Lys	Trp	Gly	Ile	Gly 710	Tyr	His	Leu	Ser	Leu 715	His	Leu	Asn	Glu	Arg 720
Cys	Asp	Pro	Glu	Ser 725	Ile	Thr	Ser	Leu	Val 730	Lys	Gln	His	Ile	Ser 735	Asp
Ala	Lys	Leu	Thr 740	Ala	Gln	Ser	Glu	Glu 745	Lys	Leu	Val	Tyr	Ile 750	Leu	Pro
Leu	Glu	Arg 755	Thr	Asn	Lys	Phe	Pro 760	Glu	Leu	Tyr	Arg	Asp 765	Leu	Asp	Arg
Cys	Ser 770	Asn	Gln	Gly	Ile	Glu 775	Asp	Tyr	Gly	Val	Ser 780	Ile	Thr	Thr	Leu
Asn 785	Glu	Val	Phe	Leu	Lys 790	Leu	Glu	Gly	Lys	Ser 795	Thr	Ile	Asp	Glu	Ser 800

Asp Ile Gly Ile Trp 805 Gly Gln Leu Gln Thr 810 Asp Gly Ala Lys Asp 815 Ile  
 Gly Ser Leu Val 820 Glu Leu Glu Gln Val 825 Leu Ser Ser Phe His 830 Glu Thr  
 Arg Lys Thr 835 Ile Ser Gly Val Ala 840 Leu Trp Arg Gln Gln 845 Val Cys Ala  
 Ile Ala 850 Lys Val Arg Phe Leu 855 Lys Leu Lys Lys Glu 860 Arg Lys Ser Leu  
 Trp 865 Thr Ile Leu Leu Leu 870 Phe Gly Ile Ser Phe 875 Ile Pro Gln Leu Leu 880  
 Glu His Leu Phe Tyr 885 Glu Ser Tyr Gln Lys 890 Ser Tyr Pro Trp Glu 895 Leu  
 Ser Pro Asn Thr 900 Tyr Phe Leu Ser Pro 905 Gly Gln Gln Pro Gln 910 Asp Pro  
 Leu Thr His 915 Leu Leu Val Ile Asn 920 Lys Thr Gly Ser Thr 925 Ile Asp Asn  
 Phe Leu 930 His Ser Leu Arg Arg 935 Gln Asn Ile Ala Ile 940 Glu Val Asp Ala  
 Phe 945 Gly Thr Arg Asn Gly 950 Thr Asp Asp Pro Ser 955 Tyr Asn Gly Ala Ile 960  
 Ile Val Ser Gly Asp 965 Glu Lys Asp His Arg 970 Phe Ser Ile Ala Cys 975 Asn  
 Thr Lys Arg Leu 980 Asn Cys Phe Pro Val 985 Leu Leu Asp Val Ile 990 Ser Asn  
 Gly Leu Leu 995 Gly Ile Phe Asn Ser 1000 Ser Glu His Ile Gln 1005 Thr Asp Arg  
 Ser Thr 1010 Phe Phe Glu Glu 1015 His Met Asp Tyr Glu Tyr 1020 Gly Tyr Arg Ser  
 Asn Thr 1025 Phe Phe Trp Ile 1030 Pro Met Ala Ala 1035 Ser Phe Thr Pro Tyr Ile 1040  
 Ala Met Ser Ser 1045 Ile Gly Asp Tyr Lys Lys 1050 Lys Ala His Ser Gln Leu 1055  
 Arg Ile Ser Gly 1060 Leu Tyr Pro Ser Ala Tyr Trp Phe Gly Gln 1070 Ala Leu  
 Val Asp Val 1075 Ser Leu Tyr Phe Leu 1080 Ile Leu Leu Leu Met 1085 Gln Ile Met



Asp Tyr Ile Phe Ser Pro Glu Glu Ile Ile Phe Ile Ile Gln Asn Leu  
 1090 1095 1100  
 Leu Ile Gln Ile Leu Cys Ser Ile Gly Tyr Val Ser Ser Leu Val Phe  
 1105 1110 1115 1120  
 Leu Thr Tyr Val Ile Ser Phe Ile Phe Arg Asn Gly Arg Lys Asn Ser  
 1125 1130 1135  
 Gly Ile Trp Ser Phe Phe Phe Leu Ile Val Val Ile Phe Ser Ile Val  
 1140 1145 1150  
 Ala Thr Asp Leu Asn Glu Tyr Gly Phe Leu Gly Leu Phe Phe Gly Thr  
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 Met Leu Ile Pro Pro Phe Thr Leu Ile Gly Ser Leu Phe Ile Phe Ser  
 1170 1175 1180  
 Glu Ile Ser Pro Asp Ser Met Asp Tyr Leu Gly Ala Ser Glu Ser Glu  
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 Ile Val Tyr Leu Ala Leu Leu Ile Pro Tyr Leu His Phe Leu Ile Phe  
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 Leu Phe Ile Leu Arg Cys Leu Glu Met Asn Cys Arg Lys Lys Leu Met  
 1220 1225 1230  
 Arg Lys Asp Pro Val Phe Arg Ile Ser Pro Arg Ser Asn Ala Ile Phe  
 1235 1240 1245  
 Pro Asn Pro Glu Glu Pro Glu Gly Glu Glu Glu Asp Ile Gln Met Glu  
 1250 1255 1260  
 Arg Met Arg Thr Val Asn Ala Met Ala Val Arg Asp Phe Asp Glu Thr  
 1265 1270 1275 1280  
 Pro Val Ile Ile Ala Ser Cys Leu Arg Lys Glu Tyr Ala Gly Lys Lys  
 1285 1290 1295  
 Lys Asn Cys Phe Ser Lys Arg Lys Lys Thr Ile Ala Thr Arg Asn Val  
 1300 1305 1310  
 Ser Phe Cys Val Lys Lys Gly Glu Val Ile Gly Leu Leu Gly His Asn  
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 Gly Ala Gly Lys Ser Thr Thr Ile Lys Met Ile Thr Gly Asp Thr Lys  
 1330 1335 1340  
 Pro Thr Ala Gly Gln Val Ile Leu Lys Gly Ser Gly Gly Gly Glu Pro  
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 33

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 <212> PRT  
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<400> 8

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			20					25					30		
His	Glu	Met	Val	Gly	Val	Ile	Phe	Ser	Asp	Thr	Phe	Ser	Tyr	Arg	Leu
		35					40					45			
Lys	Phe	Asn	Trp	Gly	Tyr	Arg	Ile	Pro	Val	Ile	Lys	Glu	His	Ser	Glu
	50					55					60				
Tyr	Thr	Glu	His	Cys	Trp	Ala	Met	His	Gly	Glu	Ile	Phe	Cys	Tyr	Leu
65					70					75					80
Ala	Lys	Tyr	Trp	Leu	Lys	Gly	Phe	Val	Ala	Phe	Gln	Ala	Ala	Ile	Asn
				85					90					95	
Ala	Ala	Ile	Ile	Glu	Val	Thr	Thr	Asn	His	Ser	Val	Met	Glu	Glu	Leu
			100					105					110		
Thr	Ser	Val	Ile	Gly	Ile	Asn	Met	Lys	Ile	Pro	Pro	Phe	Ile	Ser	Lys
		115					120					125			
Gly	Glu	Ile	Met	Asn	Glu	Trp	Phe	His	Phe	Thr	Cys	Leu	Val	Ser	Phe
	130					135					140				
Ser	Ser	Phe	Ile	Tyr	Phe	Ala	Ser	Leu	Asn	Val	Ala	Arg	Glu	Arg	Gly
145					150					155					160
Lys	Phe	Lys	Lys	Leu	Met	Thr	Val	Met	Gly	Leu	Arg	Glu	Ser	Ala	Phe
				165					170					175	
Trp	Leu	Ser	Trp	Xaa	Leu	Thr	Tyr	Ile	Cys	Phe	Ile	Phe	Ile	Met	Ser
			180					185					190		
Ile	Phe	Met	Ala	Leu	Val	Ile	Thr	Ser	Ile	Ser	Ile	Val	Phe	His	Thr
		195					200					205			
Gly	Phe	Met	Val	Ile	Phe	Thr	Leu	Tyr	Ser	Leu	Tyr	Gly	Leu	Ser	Leu
	210					215					220				
Ile	Ala	Leu	Ala	Phe	Leu	Met	Ser	Val	Leu	Ile	Arg	Lys	Pro	Met	Leu
225					230					235					240
Ala	Gly	Leu	Ala	Gly	Phe	Leu	Phe	Thr	Val	Phe	Trp	Gly	Cys	Leu	Gly
				245					250					255	

Phe Thr Val Leu Tyr Arg Gln Leu Pro Leu Ser Leu Gly Trp Val Leu  
 260 265 270  
 Ser Leu Leu Ser Pro Phe Ala Phe Thr Ala Gly Met Ala Gln Val Thr  
 275 280 285  
 His Leu Asp Asn Tyr Leu Ser Gly Val Ile Phe Pro Asp Pro Ser Gly  
 290 295 300  
 Asp Ser Tyr Lys Met Ile Ala Thr Phe Phe Ile Leu Ala Phe Asp Thr  
 305 310 315 320  
 Leu Phe Tyr Leu Ile Phe Thr Leu Tyr Phe Glu Arg Val Leu Pro Asp  
 325 330 335  
 Lys Asp Gly His Gly Asp Ser Pro Leu Phe Phe Leu Lys Ser Ser Phe  
 340 345 350  
 Trp Ser Lys His Gln Asn Thr His His Glu Ile Phe Glu Asn Glu Ile  
 355 360 365  
 Asn Pro Glu His Ser Ser Asp Asp Ser Phe Glu Pro Val Ser Pro Glu  
 370 375 380  
 Phe His Gly Lys Glu Ala Ile Arg Ile Arg Asn Val Ile Lys Glu Tyr  
 385 390 395 400  
 Asn Gly Lys Thr Gly Lys Val Glu Ala Leu Gln Gly Ile Phe Phe Asp  
 405 410 415  
 Ile Tyr Glu Gly Gln Ile Thr Ala Ile Leu Gly His Asn Gly Ala Gly  
 420 425 430  
 Lys Ser Thr Leu Leu Asn Ile Leu Ser Gly Leu Ser Val Ser Thr Glu  
 435 440 445  
 Gly Ser Ala Thr Ile Tyr Asn Thr Gln Leu Ser Glu Ile Thr Asp Met  
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 Glu Glu Ile Arg Lys Asn Ile Gly Phe Cys Pro Gln Phe Asn Phe Gln  
 465 470 475 480  
 Phe Asp Phe Leu Thr Val Arg Glu Asn Leu Arg Val Phe Ala Lys Ile  
 485 490 495  
 Lys Gly Ile Gln Pro Lys Glu Val Glu Gln Glu Val Lys Arg Ile Ile  
 500 505 510  
 Met Glu Leu Asp Met Gln Ser Ile Gln Asp Ile Ile Ala Lys Lys Leu  
 515 520 525  
 Ser Gly Gly Gln Lys Arg Lys Leu Thr Leu Gly Ile Ala Ile Leu Gly  
 530 535 540

Asp 545	Pro	Gln	Val	Leu	Leu 550	Leu	Asp	Glu	Pro	Thr 555	Ala	Gly	Leu	Asp	Pro 560
Phe	Ser	Arg	His	Arg 565	Val	Trp	Ser	Leu	Leu 570	Lys	Glu	His	Lys	Val 575	Asp
Arg	Leu	Ile	Leu 580	Phe	Ser	Thr	Gln	Phe 585	Met	Asp	Glu	Ala	Asp 590	Ile	Leu
Ala	Asp	Arg 595	Lys	Val	Phe	Leu	Ser 600	Asn	Gly	Lys	Leu	Lys 605	Cys	Ala	Gly
Ser	Ser 610	Leu	Phe	Leu	Lys	Arg 615	Lys	Trp	Gly	Ile	Gly 620	Tyr	His	Leu	Ser
Leu 625	His	Arg	Asn	Glu	Met 630	Cys	Asp	Thr	Glu	Lys 635	Ile	Thr	Ser	Leu	Ile 640
Lys	Gln	His	Ile	Pro 645	Asp	Ala	Lys	Leu	Thr 650	Thr	Glu	Ser	Glu	Glu 655	Lys
Leu	Val	Tyr	Ser 660	Leu	Pro	Leu	Glu	Lys 665	Thr	Asn	Lys	Phe	Pro 670	Asp	Leu
Tyr	Ser	Asp 675	Leu	Asp	Lys	Cys	Ser 680	Asp	Gln	Gly	Ile	Arg 685	Asn	Tyr	Ala
Val	Ser 690	Val	Thr	Ser	Leu	Asn 695	Glu	Val	Phe	Leu	Asn 700	Leu	Glu	Gly	Lys
Ser 705	Ala	Ile	Asp	Glu	Pro 710	Asp	Phe	Asp	Ile	Gly 715	Lys	Gln	Glu	Lys	Ile 720
His	Val	Thr	Arg	Asn 725	Thr	Gly	Asp	Glu	Ser 730	Glu	Met	Glu	Gln	Val 735	Leu
Cys	Ser	Leu	Pro 740	Glu	Thr	Arg	Lys	Ala 745	Val	Ser	Ser	Ala	Ala 750	Leu	Trp
Arg	Arg	Gln 755	Ile	Tyr	Ala	Val	Ala 760	Thr	Leu	Arg	Phe	Leu 765	Lys	Leu	Arg
Arg	Glu 770	Arg	Arg	Ala	Leu	Leu 775	Cys	Leu	Leu	Leu	Val 780	Leu	Gly	Ile	Ala
Phe 785	Ile	Pro	Ile	Ile	Leu 790	Glu	Lys	Ile	Met	Tyr 795	Lys	Val	Thr	Arg	Glu 800
Thr	His	Cys	Trp	Glu 805	Phe	Ser	Pro	Ser	Met 810	Tyr	Phe	Leu	Ser	Leu 815	Glu
Gln	Ile	Pro	Lys	Thr	Pro	Leu	Thr	Ser	Leu	Leu	Ile	Val	Asn	Asn	Thr

820										825					830				
Gly	Ser	Asn	Ile	Glu	Asp	Leu	Val	His	Ser	Leu	Lys	Cys	Gln	Asp	Ile				
		835					840					845							
Val	Leu	Glu	Ile	Asp	Asp	Phe	Arg	Asn	Arg	Asn	Gly	Ser	Asp	Asp	Pro				
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Ser	Tyr	Asn	Gly	Ala	Ile	Ile	Val	Ser	Gly	Asp	Gln	Lys	Asp	Tyr	Arg				
865					870					875					880				
Phe	Ser	Val	Ala	Cys	Asn	Thr	Lys	Lys	Leu	Asn	Cys	Phe	Pro	Val	Leu				
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Met	Gly	Ile	Val	Ser	Asn	Ala	Leu	Met	Gly	Ile	Phe	Asn	Phe	Thr	Glu				
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Leu	Ile	Gln	Thr	Glu	Ser	Thr	Ser	Phe	Ser	Arg	Asp	Asp	Ile	Val	Leu				
		915					920					925							
Asp	Leu	Gly	Phe	Ile	Asp	Gly	Ser	Ile	Phe	Leu	Leu	Leu	Ile	Thr	Asn				
	930					935					940								
Cys	Val	Ser	Pro	Phe	Ile	Gly	Met	Ser	Ser	Ile	Ser	Asp	Tyr	Lys	Lys				
945					950					955					960				
Asn	Val	Gln	Ser	Gln	Leu	Trp	Ile	Ser	Gly	Leu	Trp	Pro	Ser	Ala	Tyr				
				965					970					975					
Trp	Cys	Gly	Gln	Ala	Leu	Val	Asp	Ile	Pro	Leu	Tyr	Phe	Leu	Ile	Leu				
			980					985					990						
Phe	Ser	Ile	His	Leu	Ile	Tyr	Tyr	Phe	Ile	Phe	Leu	Gly	Phe	Gln	Leu				
		995				1000					1005								
Ser	Trp	Glu	Leu	Met	Phe	Val	Leu	Val	Val	Cys	Ile	Ile	Gly	Cys	Ala				
	1010					1015				1020									
Val	Ser	Leu	Ile	Phe	Leu	Thr	Tyr	Val	Leu	Ser	Phe	Ile	Phe	Arg	Lys				
1025					1030				1035					1040					
Trp	Arg	Lys	Asn	Asn	Gly	Phe	Trp	Ser	Phe	Gly	Phe	Phe	Ile	Ile	Leu				
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Ile	Cys	Val	Ser	Thr	Ile	Met	Val	Ser	Thr	Gln	Tyr	Glu	Lys	Leu	Asn				
		1060					1065					1070							
Leu	Ile	Leu	Cys	Met	Ile	Phe	Ile	Pro	Ser	Phe	Thr	Leu	Leu	Gly	Tyr				
	1075					1080					1085								
Val	Met	Leu	Leu	Ile	Gln	Leu	Asp	Phe	Met	Arg	Asn	Leu	Asp	Ser	Leu				
	1090				1095					1100									

Asp Asn Arg Ile Asn Glu Val Asn Lys Thr Ile Leu Leu Thr Thr Leu  
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 Ile Pro Tyr Leu Gln Ser Val Ile Phe Leu Phe Val Ile Arg Cys Leu  
 1125 1130 1135  
 Glu Met Lys Tyr Gly Asn Glu Ile Met Asn Lys Asp Pro Val Phe Arg  
 1140 1145 1150  
 Ile Ser Pro Arg Ser Arg Glu Thr His Pro Asn Pro Glu Glu Pro Glu  
 1155 1160 1165  
 Glu Glu Asp Glu Asp Val Gln Ala Glu Arg Val Gln Ala Ala Asn Ala  
 1170 1175 1180  
 Leu Thr Ala Pro Asn Leu Glu Glu Glu Pro Val Ile Thr Ala Ser Cys  
 1185 1190 1195 1200  
 Leu His Lys Glu Tyr Tyr Glu Thr Lys Lys Ser Cys Phe Ser Thr Arg  
 1205 1210 1215  
 Lys Lys Lys Ile Ala Ile Arg Asn Val Ser Phe Cys Val Lys Lys Gly  
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 Glu Val Leu Gly Leu Leu Gly His Asn Gly Ala Gly Lys Ser Thr Ser  
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 Ile Lys Met Ile Thr Gly Cys Thr Lys Pro Thr Ala Gly Val Val Val  
 1250 1255 1260  
 Leu Gln Gly Ser Arg Ala Ser Val Arg Gln Gln His Asp Asn Ser Leu  
 1265 1270 1275 1280  
 Lys Phe Leu Gly Tyr Cys Pro Gln Glu Asn Ser Leu Trp Pro Lys Leu  
 1285 1290 1295  
 Thr Met Lys Glu His Leu Glu Leu Tyr Ala Ala Val Lys Gly Leu Gly  
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 Lys Glu Asp Ala Ala Leu Ser Ile Ser Arg Leu Val Glu Ala Leu Lys  
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 Leu Gln Glu Gln Leu Lys Ala Pro Val Lys Thr Leu Ser Glu Gly Ile  
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 Lys Arg Lys Leu Cys Phe Val Leu Ser Ile Leu Gly Asn Pro Ser Val  
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 Val Leu Leu Asp Glu Pro Phe Thr Gly Met Asp Pro Glu Gly Gln Gln  
 1365 1370 1375  
 Gln Met Trp Gln Ile Leu Gln Ala Thr Val Lys Asn Lys Glu Arg Gly  
 1380 1385 1390  
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1. The first part of the report, "Introduction", discusses the importance of the study and the objectives of the research. It also mentions the scope of the study and the limitations of the data.

2. The second part, "Literature Review", provides a comprehensive overview of the existing literature on the topic. It covers various aspects of the research, including theoretical frameworks, empirical studies, and policy implications.

3. The third part, "Methodology", describes the research design and the methods used for data collection and analysis. It includes details about the sample size, data sources, and statistical techniques employed.

4. The fourth part, "Results", presents the findings of the study. It includes tables and figures that illustrate the data and the statistical results. The results are discussed in the context of the research objectives and the existing literature.

5. The fifth part, "Conclusion", summarizes the main findings of the study and provides recommendations for future research. It also discusses the implications of the findings for policy and practice.

6. The final part, "References", lists the sources of the literature cited in the report. It includes books, journal articles, and other relevant documents.

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